

Poster pitches
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TRICHOCOCCUS SPECIES AS CATALYSTS FOR BIOTECHNOLOGICAL PRODUCTION OF 1,3-PROPANEDIOL

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Nowadays, large amounts of waste glycerol are generated as a major byproduct from biodiesel industry. Glycerol can be converted to 1,3-propanediol (1,3-PDO), a building block for the synthesis of polyester and biodegradable plastic fibers. We have previously isolated a novel *Trichococcus* species, strain ES5, which is capable to anaerobically produce 1,3-PDO from glycerol, an ability it shares with *Trichococcus pasteurii*. In this work, we analyzed and compared the genomes of these two bacteria and studied the production of 1,3-PDO by cultures of *T. pasteurii* and strain ES5 under different conditions. A continuous bioreactor system was established for achieving production of 1,3-PDO. A complete operon structure composed by 16 genes related to 1,3-PDO metabolism could be identified in the genomes of *T. pasteurii* and strain ES5. This genomic synteny contains the two essential genes for glycerol conversion to 1,3-PDO, glycerol dehydratase and 1,3-propanediol dehydrogenase, and additional genes for glycerol uptake and regulation factors of glycerol metabolism. The metabolic trait of 1,3-production was *in silico* analysed for the complete bacterial kingdom and genes were identified with an essential or accessory influence for the trait.

Bacteria with fewer genes were underperforming when tested *in vitro* for production 1,3-PDO. Furthermore, both *Trichococcus* strains could grow on glycerol in a broad range of temperatures, from 4 to 40°C and strain ES5 achieved growth in 0°C. Such physiological properties may be advantageous for biotechnological production. Tolerance to high salinity may be an additional strength of *Trichococcus* species as a candidate for the biotechnological production of 1,3-PDO.

Keywords

glycerol; fermentation; 1,3-propanediol; low temperature; high-salinity; genome comparison